

SEQUENCE LISTING

<110> Hagen, Gustav
 Siegmund, Hans-Ulrich
 Weichel, Walter
 Wick, Maresa
 Zubov, Dmitry

<120> Human Catalytic Telomerase Sub-Unit and its Diagnostic and Therapeutic Use

TECH CENTER 1600/2900

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<140> US 09/424,686

<141> 1999-11-29

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<212> DNA
<213> Human

<221> CDS

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- <210> 9
- <211> 4006
- <212> DNA
- <213> Human
- <220>
- <221> CDS
- <222> (1)..(4006)
- <223> Nucleotides 2184 to 2219 of SEQ ID NO. 1 have been deleted to provide this sequence.

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<211> 3824

<212> DNA ***

<213> Human

<220>

<221> CDS

<222> (1)..(3824)

<223> Nucleotides 2184 to 2219 and 2345 to 2526 of SEQ ID NO. 1 were de leted.

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<210> 11

<211> 3411 PK

<212> DNA

<213> Human

<220>

<221> CDS

<222> (1)..(3411)

<223> Nucleotides 1-59 and 3471-4042 of SEQ ID NO 1 were deleted to provide this sequence.

<400> 11 gcgatgccgc gcgctccccg ctgccgagcc gtgcgctccc tgctgcgcag ccactaccgc 60 gaggtgctgc.cgctggccac gttcgtgcgg cgcctggggc cccagggctg gcggctggtg 120 180 cagegegggg acceggegge ttteegegeg etggtggeec agtgeetggt gtgegtgeec 240 tgggacgcac ggccgccccc cgccgccccc tccttccgcc aggtgtcctg cctgaaggag ctggtggccc gagtgctgca gaggctgtgc gagcgcggcg cgaagaacgt gctggccttc 300 360 ggettegege tgetggaegg ggeeegeggg ggeeeceeg aggeetteae caccagegtg cgcagctacc tgcccaacac ggtgaccgac gcactgcggg ggagcggggc gtggggctg 420 ctgctgcgcc gcgtgggcga cgacgtgctg gttcacctgc tggcacgctg cgcgctcttt 480 gtgctggtgg ctcccagctg cgcctaccag gtgtgcgggc cgccgctgta ccagctcggc 540 gctgccactc aggcccggcc cccgccacac gctagtggac cccgaaggcg tctgggatgc 600 gaacgggcct ggaaccatag cgtcagggag gccggggtcc ccctgggcct gccagccccg 660 ggtgcgagga ggcgcggggg cagtgccagc cgaagtctgc cgttgcccaa gaggcccagg 720 cgtggcgctg cccctgagcc ggagcggacg cccgttgggc aggggtcctg ggcccacccg 780 ggcaggacgc gtggaccgag tgaccgtggt ttctgtgtgg tgtcacctgc cagacccgcc 840

900 gaagaagcca cctctttgga gggtgcgctc tctggcacgc gccactccca cccatccgtg 960 ggccgccagc accacgcggg ccccccatcc acatcgcggc caccacgtcc ctgggacacg 1020 cettqteece eggtgtaege egagaceaag caetteetet acteeteagg egacaaggag 1080 cagetgegge cetectteet acteagetet etgaggeeca geetgaetgg egeteggagg 1140 ctcgtggaga ccatctttct gggttccagg ccctggatgc cagggactcc ccgcaggttg ccccgcctgc cccagcgcta ctggcaaatg cggcccctgt ttctggagct gcttgggaac 1200 1260 cacgcgcagt gcccctacgg ggtgctcctc aagacgcact gcccgctgcg agctgcggtc accccagcag coggtgtctg tgcccgggag aagccccagg gctctgtggc ggcccccgag 1320 1380 gaggaggaca cagacccccg tcgcctggtg cagctgctcc gccagcacag cagcccctgg caggtgtacg gcttcgtgcg ggcctgcctg cgccggctgg tgcccccagg cctctggggc 1440 tccaggcaca acgaacgccg cttcctcagg aacaccaaga agttcatctc cctggggaag 1500 catgccaagc tctcgctgca ggagctgacg tggaagatga gcgtgcggga ctgcgcttgg 1560 ctgcgcagga gcccaggggt tggctgtgtt ccggccgcag agcaccgtct gcgtgaggag 1620 1680 atcctggcca agttcctgca ctggctgatg agtgtgtacg tcgtcgagct gctcaggtct 1740 ttcttttatg tcacggagac cacgtttcaa aagaacaggc tctttttcta ccggaagagt gtctggagca agttgcaaag cattggaatc agacagcact tgaagagggt gcagctgcgg 1800 gagetgtegg aageagaggt caggeageat egggaageea ggeeegeeet getgaegtee 1860 agactccgct tcatccccaa gcctgacggg ctgcggccga ttgtgaacat ggactacgtc 1920 gtgggagcca gaacgttccg cagagaaaag agggccgagc gtctcacctc gagggtgaag 1980 gcactgttca gcgtgctcaa ctacgagcgg gcgcggcgcc ccggcctcct gggcgcctct 2040 gtgctgggcc tggacgatat ccacagggcc tggcgcacct tcgtgctgcg tgtgcgggcc 2100 caggacccgc cgcctgagct gtactttgtc aaggtggatg tgacgggcgc gtacgacacc 2160 atcccccagg acaggctcac ggaggtcatc gccagcatca tcaaacccca gaacacgtac 2220 2280 tgcgtgcgtc ggtatgccgt ggtccagaag gccgcccatg ggcacgtccg caaggccttc aagagccacg tototacott gacagacoto cagoogtaca tgcgacagtt cgtggctcac 2340 ctgcaggaga ccagcccgct gagggatgcc gtcgtcatcg agcagagctc ctccctgaat 2400 2460 gaggccagca gtggcctctt cgacgtcttc ctacgcttca tgtgccacca cgccgtgcgc atcaggggca agtcctacgt ccagtgccag gggatcccgc agggctccat cctctccacg 2520 ctgctctgca gcctgtgcta cggcgacatg gagaacaagc tgtttgcggg gattcggcgg 2580 gacgggctgc tcctgcgttt ggtggatgat ttcttgttgg tgacacctca cctcacccac 2640 gcgaaaacct tcctcaggac cctggtccga ggtgtccctg agtatggctg cgtggtgaac 2700 ttgcggaaga cagtggtgaa cttccctgta gaagacgagg ccctgggtgg cacggctttt 2760 gttcagatgc cggcccacgg cctattcccc tggtgcggcc tgctgctgga tacccggacc 2820 ctggaggtgc agagcgacta ctccagctat gcccggacct ccatcagagc cagtctcacc 2880 ttcaaccgcg gcttcaaggc tgggaggaac atgcgtcgca aactctttgg ggtcttgcgg 2940 3000 ctgaagtgtc acagcctgtt tctggatttg caggtgaaca gcctccagac ggtgtgcacc aacatctaca agatcctcct gctgcaggcg tacaggtttc acgcatgtgt gctgcagctc 3060 ccatttcatc agcaagtttg gaagaacccc acatttttcc tgcgcgtcat ctctgacacg 3120 gcctccctct gctactccat cctgaaagcc aagaacgcag ggatgtcgct gggggccaag 3180 ggcgccgccg gccctctgcc ctccgaggcc gtgcagtggc tgtgccacca agcattcctg 3240 ctcaagctga ctcgacaccg tgtcacctac gtgccactcc tggggtcact caggacagcc 3300 cagacgcagc tgagtcggaa gctcccgggg acgacgctga ctgccctgga ggccgcagcc 3360 3411 aacceggeac tgeecteaga etteaagace ateetggaet gatggeeace e

<210> 12

<211> 4012

<212> DNA

<213> Homo sapien

<220>

<221> CDS

<222> (1)..(4042)

Nucleotide positions 1-1782 and 3872 to 4042 are identical to the same sequences in SEQ ID NO: 1; nucleotide positions from 1783 to 3871 are according to SEQ ID NO: 7.

<400> 12

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gactccgctt catccccaag cctgacgggc tgcggccgat tgtgaacatg gactacgtcg 1980 2040 tgggagccag aacgttccgc agagaaaaga gggccgagcg tctcacctcg agggtgaagg cactgttcag cgtgctcaac tacgagcggg cgcggcgccc cggcctcctg ggcgcctctg 2100 2160 tgctgggcct ggacgatatc cacagggcct ggcgcacctt cgtgctgcgt gtgcgggccc aggacccgcc gcctgagctg tactttgtca aggtggatgt gacgggcgcg tacgacacca 2220 2280 tcccccagga caggctcacg gaggtcatcg ccagcatcat caaaccccag aacacgtact 2340 gegtgegteg gtatgeegtg gtecagaagg cegeceatgg geaegteege aaggeettea agagecacgt ctctacettg acagacetee agecgtacat gegacagtte gtggeteace 2400 tgcaggagac cagcccgctg agggatgccg tcgtcatcga gcagagctcc tccctgaatg 2460 2520 aggccagcag tggcctcttc gacgtcttcc tacgcttcat gtgccaccac gccgtgcgca 2580 tcaggggcaa gtcctacgtc cagtgccagg ggatcccgca gggctccatc ctctccacgc 2640 tgctctgcag cctgtgctac ggcgacatgg agaacaagct gtttgcgggg attcggcggg acgggctgct cctgcgtttg gtggatgatt tcttgttggt gacacctcac ctcacccacg 2700 cgaaaacctt cctcaggacc ctggtccgag gtgtccctga gtatggctgc gtggtgaact 2760 tgcggaagac agtggtgaac ttccctgtag aagacgaggc cctgggtggc acggcttttg 2820 ttcagatgcc ggcccacggc ctattcccct ggtgcggcct gctgctggat acccggaccc 2880 tggaggtgca gagcgactac tccagctatg cccggacctc catcagagcc agtctcacct 2940 3000 tcaaccgcgg cttcaaggct gggaggaaca tgcgtcgcaa actctttggg gtcttgcggc 3060 tgaagtgtca cagcctgttt ctggatttgc aggtgaacag cctccagacg gtgtgcacca acatetacaa gateeteetg etgeaggegt acaggtttea egeatgtgtg etgeagetee 3120 catttcatca gcaagtttgg aagaacccca catttttcct gcgcgtcatc tctgacacgg 3180 cctcctctg ctactccatc ctgaaagcca agaacgcagg tatgtgcagg tgcctggcct 3240 cagtggcagc agtgcctgcc tgctggtgtt agtgtgtcag gagactgagt gaatctgggc 3300 ttaggaagtt cttacccctt ttcgcatcag gaagtggttt aacccaacca ctgtcaggct 3360 3420 cgtctgcccg ccctctcgtg gggtgagcag agcacctgat ggaagggaca ggagctgtct gggagctgcc atcettccca cettgctctg cetggggaag cgctgggggg cetggtctct 3480 cctgtttgcc ccatggtggg atttgggggg cctggcctct cctgtttgcc ctgtggtggg 3540 3600 attgggctgt ctcccgtcca tggcacttag ggcccttgtg caaacccagg ccaagggctt aggaggaggc caggcccagg ctaccccacc cctctcagga gcagaggccg cgtatcacca 3660

N + 9

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